

## AMENDMENTS

### *In the specification*

On page 1, in the paragraph beginning on line 9, please amend as follows:

B1  
This application is a continuation of U.S. Serial No. 08/752,844, filed November 21, 1996, now U.S. Patent No. 5,935,821, which is a continuation-in-part of U.S. Serial No. 08/591,196, filed January 16, 1996, now U.S. Patent No. 5,977,316, which is a continuation-in-part of U.S. Serial No. 08/372,676, filed January 17, 1995, now U.S. Patent No. 5,612,030, all of which are hereby incorporated herein in their entirety.

On pages 85 and 86, in the paragraphs from line 4 on page 85 to line 5 on page 86, containing the paragraphs previously amended in the Preliminary Amendment of April 15, 1999, please amend the specification as follows:

Amongst the 50 database sequences matched most closely to that of the 1A7 heavy chain variable region, none was identical. The following summarizes the main points deduced from the comparison.

- B2
- The closest match was with a heavy chain fragment beginning at residue 9 (designation gp|M36221|MUSIGHAEB\_1). There were 6 substitutions between residues 1 and 97 (before the VDJ junction), 6 substitution differences after residue 97, and 1A7 was shorter about the VDJ junction by 2 residues.
  - The closest match with a full length heavy chain variable region had the following features (designation gp|U01185|MMU01185): There were 10 substitution differences between residues 1 and 97, 6 substitutions after residue 97, and 1A7 was shorter about the VDJ junction by 3 residues.
  - 1A7 differed in length from all sequences but one, due to insertions or deletions of 1 to 8 residues about the VDJ junction. For the sequence of equal length (designation